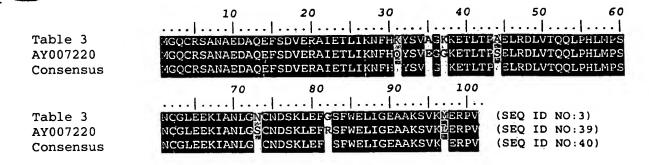
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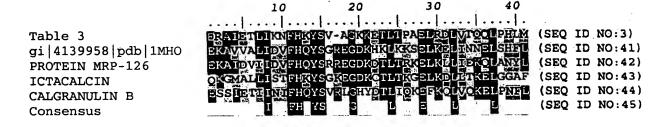
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### Figure 4C

	10	20	30	40	. <b>50</b>	60
				1		1
Table 6	MGQCRSANAEDAQEF					
AY007220	MGQCRSANAEDAQEF					
Consensus	MGQCRSANAEDAQEF	SDVERAIETI	LIKNFHQYSV	EGGKETLTPS	ELRDLVTQQL	PHLMPS
	70	80	90	100		
Table 6 AY007220 Consensus	NCGLEEKIANLGSCN	DSKLEFRSFV	ELIGEAAKS	VKLERPVRGH	(SEQ ID N	0:6)
	NCGLEEKIANLGSCN				(SEQ ID N	0:39)
	NCGLEEKIANLGSCN	DSKLEFRSF	VELIGEAAKS	VKLERPVRGH	(SEQ ID N	0:40)

## Figure 4D



### Figure 4E

30 20 10 Table 6 (SEQ ID NO:6) gi | 4139958 | pdb | 1MHO (SEQ ID NO:41) FHQYSRREGDKDTL PROTEIN MRP-126 (SEQ ID NO:42) TIINTFHOYSVELSHYDTLIGKEF: OLV (SEQ ID NO:44) CALGRANULIN B (SEQ ID NO:46) CALGRANULIN B (SEQ ID NO:47) Consensus



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# Figure 7

# 37) .. 02 CTGGTGAAAGTTCTTGATGAGGGTCTCAATGGCCCTCTCCACATCACTGAATTC (SEQ ID GAATTCCAGAGGGAGTTCTCAGTGCCCCCGGACAGGCCTCTCCAGCTTCACACTCTTGGC TGCCCCCGGACAGTCCTCTCNAGCTTCACACTCTTGGC GAATTCCAGAGGGAGTTCTCAGTGCCCCCGGACAGGCCTCTCCAGCTTCACACTCTTGGC CGCTTCTCCAATCAGCTCCCAGAAACTCCTGAACTCCAGTTTAGAGTCATTGCAGCTGCC CGCTTCTCCAATCAGCTCCCAGAAACTCCTGAACTCCAGTTTAGAGTCATTGCAGCTGCC CGCTTCTCCAATCAGCTCCCAGAAACTCCTGAACTCCAGTTTAAGAGTCATTGCAGCTGCC CAGGTTGGCAATTTTCTCTTCCAGGCCACAGTTGCTCGGCATGAGATGGGGCAGCTGCTG CAGGTTGGCAATTTTCTCTTCCAGGCCANAGTTGCTCGGCATGAGATGGGGCAGCTGCTG CAGGTTGGCAATTTTCTCTTCCAGGCCACAGTTGCTCGGCATGAGATGGGGCAGCTGCTG GGTGACCAGGTCCCGTAGCTCAGAAGGGGTCAGCGTCTCCCTTCCCACCCTCCACGGAGTA GGTGACCAGGTCCCGTAGCTCAGAAGGGGTCAGCGTCTCCTTCCCACCCTCCACGGAGTA GGTGACCAGGTCCCGTAGCTCAGAAGGGGTCAGCGTCTCCCTTCCCACCCTCCACGGAGTA CTGGTGAAAGTTCTTGATGAGGGTCTCAATGGCCCTCTCCACATCACTGAATTCCTGAGC CTGGTGAAAGTTCTTGATGAGGGTCTCAATGGCCCTCTCCACATCACTGAATTCCTGAGC ATCCTCTGCGTTGGCTGACCGACACTGTCCCATGGTGCTCACTGTGTCTGGTCCTTTTGGT ATCCTCTGCGTTGGCTGACCGACACTGTCCCATGGTGCTCACTGTGTCTGGTCCTTTTGGT GAGAGTTCTGTTGTCCTAT (SEQ ID NO: 49) 5 GAGAGTTCTGTTGTCCTAT (SEQ ID NO: consensus 65677221+ 65677221+ 65677221+ consensus consensus consensus 65677221+ consensus AA315020-65677221+ AA315020-AA315020-AA315020-AA315020-AA315020consensus AA315020consensus